FIGURE 1

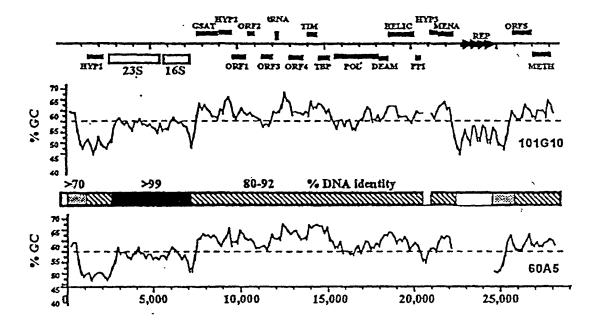


Figure 2

69.	<u>Gene</u>	<u>Strain</u>		TATA Box			Coding Start		TATA to Start (bp)	
81 82	Hypoth 03	A B					GCGGCGCATG CCGGCGCGTG			25
93 94	Hypoth 02	A B	GGAAACTTTG	ATTATA	CGGG	CGTACATTCC	CGGGGCCCAT CGGGGCCCAT	G	~~~~~~	26
85 86	ORF 02	A B	ACGGCAAGGT	AATAAT	AGCC	TGCCGTCCGT	AACGGCCGTA ACCTGCCGTA	TG	~~~~~~	27
87 88		A B	CATGGAACTA	GATAAT	AACC	GGTCCCGCGG	ATCCCATGCA GTACAATGCA	TG	~~~~~~~	27
89 9 o	regit to get the second of the	A B	AGCACGACAA	GTTATA	GCAG	GGTACAAAGG	GTGCGCGCGC AGCAGCGCAC	ATG	~~~~~~	28
92		A B	ATCCGGCCTC	ATTAAA	TTAC	GGGGGTACA	GCCTGCTGCC ACCTGCTGCC	GTG~~~~~	~~~~~~	28
94		A B	ACTTCATACA	CATAAA	TCCC	GCCTGAACGG	GCGGCTGCGC TCGTCCGCGC CACCATGGCC	ATG	~~~~~~~	29
96	deaminase	A B A	CCGCATATAC	CATAAT	ATGC	CGGGCGGGG	CAGGCTGCCC CAGGGCCGCG	.GTG~~~~~	~~~~~~~	29
98	7-107 - Ny - Ny - Ny - Ny - Ny - Ny - Ny - Ny	B A	GGGTAGAAAC	CATAAA	ACAA	CAGGCCGCGG	CAGGGCG.CG GCGCGTATCA	CGTG~~~~~	~~~~~~~	29
100	ORF 06 tRNA-tyr	B A	ATACACGTGG	TATAAA	CAGA	GG.CCGGACG	GCGCGGACCA CACGGATCGT	CATG~~~~	~~~~~~~	29
103		B A	CCGGGCCCCG	GTTAAA	ATAG	CG.CACGGGC	CACCCGTCGT GGATCCTGAC	CAATG~~~~	~~~~~~	30
104	TIM	B A	GCGTCGATAG	AATAAA	TACG	CGCAGGGGGC	GGCACCGGAT CCCGTGGCGC	GATCGCCCGT	G~~~~~	36
107		B A B	ATTTCAACTA	CATAAA	TGCC	TAGTTACGCA	GCGGTGC GAAATAGCAA GAAATATCAA	ACGACGTACT	TCGACTAATG	45
109	ORF 01	A B	ACGGCAGGCT	ATTATT	ACCT	TGCCTTGCGT	TGTA //G	CGGGGTGCGG	CAGGGGATG	52
111	Methylase	-	CTACAACGAT	TTTAAG	TCGG	CGCCGGGGCA	GCCG.//G GCGG.//T	ATGTGGGGCA	GGCAACATG	104
	16S RNA						CCGATCCGAT GCGATCCGAT			220
•	Archaeal promoter consensus			YTTAWA						

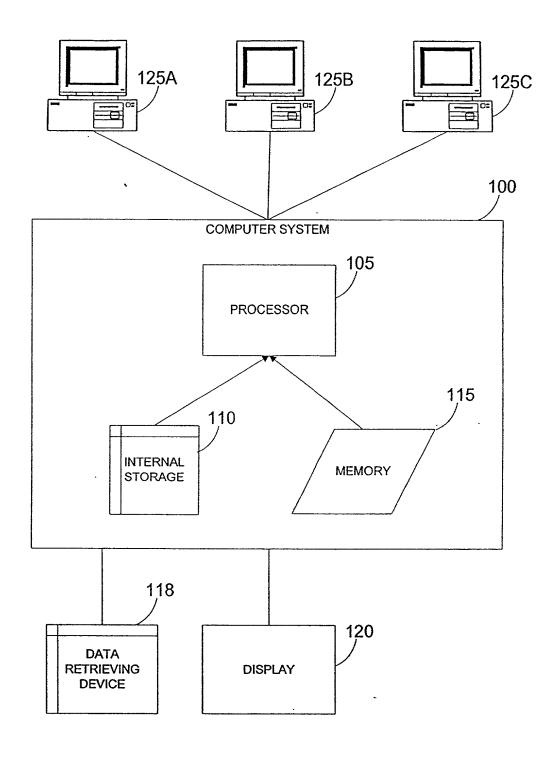


FIGURE 3

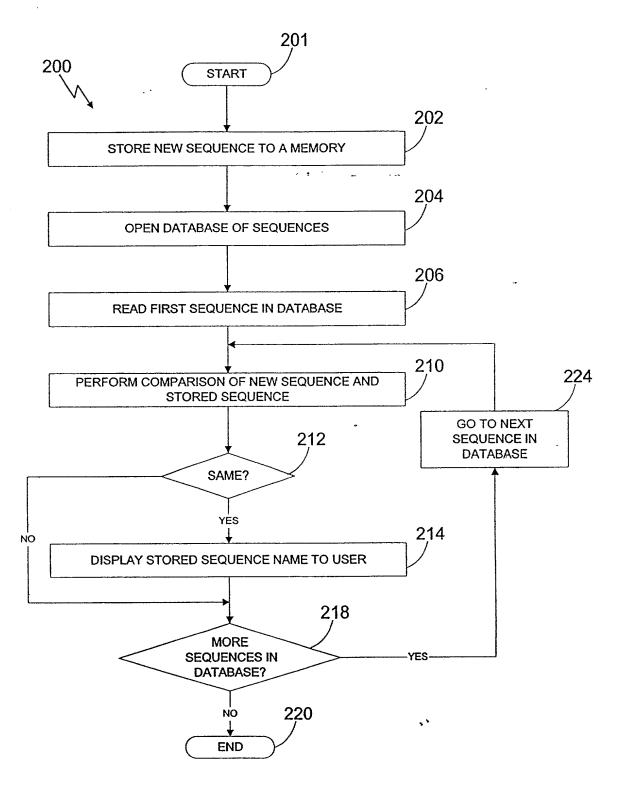


FIGURE 4

